

Fig.1

10 20 30 40 50 60
AACGGATCTG CCCGCCTCAG CCTCCCAAAG TGCTGGGATT GCAGGCGTGA GCCACCTCAC

70 80 90 100 110 120
CTGGCTACAA GTTTTCAAAA TACATTTATC TAGTACCCAT ACATTCTCCA GTTTGTCCAC

130 140 150 160 170 180
AGGACATCTT ATGACTTGAG CAAGCTGCTA AAAATCCAAG GGTGCAGCGT TTGTATGTCT

190 200 210 220 230 240
ATAGGATTGC TCAGATCTGC CCCACCCCTG AAAGAATTTA AGAGAATTTC TTGAGGCCAG

250 260 270 280 290 300
GCACAGTGGC TCACACCTGT AATTCCAGTA CTGTGAGAGT CCGAGGTCAG AGGACTGCTT

PPRE

310 320 330 340 350 360
GAGGCCAGGA GTTCAAGAGC AGCCTGGACA ACACAGGGAG ACCTGTCACT ACAAAGAATA

370 380 390 400 410 420
AATAAATTAG CCAGGCTTAG TGGCTCATCC CTGTGGTCCC AGCTACTAGG GAGGCAGAAG

430 440 450 460 470 480
TAGGACTGCT TGTCCCAGGA GGTCAAGACT GCAGTGAGCT GAGACCCAGC CACCTGCATT

490 500 510 520 530 540

CCAGCCTGGG CAACAAAAAG AGACCCTGTC TCAAAAAATA AGTTAAATAA ATAAATAATA

550 560 570 580 590 600

AAAATAGTTT AAACCCTAAA CACATCTTCT TTTTCAAAGA GGA CTCTTA AGGACTTCAT

610 620 630 640 650 660
GCTGCGTCCT GTTGATCTCC ACTTCCCTTT TTCAGCGTCC AACTTTTAA CAGTCTCTTT

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Fig.2

670 680 690 700 710 720
 TGCCAAGGAT AATAAGTATA TAGTTTCTGG AATCCAGATT CTTCCCTGTT TGGACAGCCA

 730 740 750 760 770 780
 GGGGGACAAT TTTTGGTCTG CAGGCCTTTG CATCTGTTCT GCTGTTGCTC AGCAATCTCA
 GRE
 790 800 810 820 830 840
 CAGCAAATTT GCCGAGCCTC TCCGGAATGC ACAGCCAGAC AGAGCTCAGC GCAAAAGCTA

 850 860 870 880 890 900
 GAGAACCTGG CGGAGGGAGA CTCACAGTGC CACAAAAAAA CTTTATCTTT TCTTTTTTTT

 910 920 930 940 950 960
 TTTCTTTTCT TTCTTTCTCT TTCTTTCTTG TCTTTCTGTC TTTCCTCTCT CTCTCTCTGT

 970 980 990 1000 1010 1020
 CTTTCTTTCC TCTCTTTCTT TCTTTTTTCC TACATGGCAA GATCTCCTCA TGGCAGAAAT

 1030 1040 1050 1060 1070 1080
AATCTGCCTT GACTTCTGTT TCCACGCTGC TTCTGCCAGG ACCATGCGCT CGGCGTGTTT
 GRE
 1090 1100 1110 1120 1130 1140
 TTCTTTCCGC TATAATTATC CAGGCCCATC CCAGCTCTGG TCCCCTCAGC TGTTCCCTGG

 1150 1160 1170 1180 1190 1200
 CAGTCCCTTC TGCTGGTGAA AACACATATG GCGCCGGCCT GACCAGGGTG TAAGTGTGTG

 1210 1220 1230 1240 1250 1260
 AATATCAGGA AGATGACTGA ACGTCTTTGG GACTCCGTTT CCTCATTGTA AAATGGAGGT

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[illegible]

1270 1280 1290 1300 1310 1320
TAATACCAGC CTTCTTCTAC TCCCCAAACG CACGTGTTTG TCCCGGCCAG AGGGCCCAAT
C/EBP

1330 1340 1350 1360 1370 1380
TGTTGGCTGT TCACGCATCA GTTACCCCCA CAGGACGGGT CAGCCAATTA AAGGCGAACC
C/EBP

1390 1400 1410 1420 1430 1440
AGGCCCCGGTC CATCTCCTGA CGCCTTTTCT CATCCCAGGG CTGGACAGGC AGCTGGCCTG
MyoD

1450 1460 1470 1480 1490 1500
GGCCCCGGCTC TGCCITGTCA CGTGCGGGGG CCGGCCCGTT TGCTTGTTCTG TGTGTAGGAG
GRE

1510 1520 1530 1540 1550 1560
CGTGAGGTCA CGCTGGGTGC TCCCGCCCCG CCGGGGCCCTT TAGTGTCCCT GGTCCCTAAA

1570 1580 1590 1600 1610 1620
CGCCAGGCCG CTCCACCGGG GGAGAAGGCG CGAACCCCAG CCGAGCCCAA CGGCTGTTGT

1630 1640 1650 1660 1670 1680
CGGTTGCCGG GCCACCTGTT GCTGCAGTTC TGATTGGTTC CTCCCCCGA CAACGCGGCG

1690 1700 1710 1720 1730 1740
GCTGTAACCA ATCGACAGCG AGGCCGGTCG CGAGGCCCCA GTCCCGCCCT GCAGGAGCCA
C/EBP

1750 1760 1770 1780 1790 1800
GCCGCGCGCT CGCTCGCAGG AGGGTGGGTA GTTTGCCCAG CGTAGGGGGG CTGGGCCCCAT

1810 1820 1830 1840 1850 1860
AAAAGAGGAA GTGCACTTAA GACACGGCCC CGCTGGACGC TTGTTAGAAA CCGTCCTGGC

1870 1880 1890 1900 1910 1920
TGGAAGGCA AGAGGTGTGT GACTGGACAA GACTTGTTTC TGGCGGTCAG TCTTGCCATC